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Sequence Listing was accepted.

See attached Validation Report.

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Reviewer: Keisha Douglas

Timestamp: Tue Aug 28 14:32:26 EDT 2007

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Application No: 10568414 Version No: 2.1

**Input Set:**

**Output Set:**

**Started:** 2007-08-28 14:31:20.508  
**Finished:** 2007-08-28 14:31:21.124  
**Elapsed:** 0 hr(s) 0 min(s) 0 sec(s) 616 ms  
**Total Warnings:** 0  
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**No. of SeqIDs Defined:** 2  
**Actual SeqID Count:** 2

SEQUENCE LISTING

<110> Aasly, Jan O  
Wszolek, Zbigniew K  
Farrer, Matthew J

<120> Polynucleotide

<130> U1913

<140> 10/568,414  
<141> 2006-02-14

<150> NO20045612  
<151> 2004-12-23

<160> 2

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<222> (2019)..(2019)  
<223> X may be any amino acid except glycine.

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<309> 2004-11-15  
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35 40 45

Glu His Ala Ser Lys Leu Phe Gln Gly Lys Asn Ile His Val Pro Leu  
50 55 60

Leu Ile Val Leu Asp Ser Tyr Met Arg Val Ala Ser Val Gln Gln Val  
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Gly Trp Ser Leu Leu Cys Lys Leu Ile Glu Val Cys Pro Gly Thr Met  
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Gln Ser Leu Met Gly Pro Gln Asp Val Gly Asn Asp Trp Glu Val Leu  
100 105 110

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115 120 125

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Ser Gly Lys Ile Thr Leu Leu Ile Leu Asp Glu Glu Ser Asp Ile Phe  
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Met Leu Ile Phe Asp Ala Met His Ser Phe Pro Ala Asn Asp Glu Val  
165 170 175

Gln Lys Leu Gly Cys Lys Ala Leu His Val Leu Phe Glu Arg Val Ser  
180 185 190

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195 200 205

Leu Ser Ala Ser Thr Asn Phe Lys Asp Glu Glu Glu Ile Val Leu His  
210 215 220

Val Leu His Cys Leu His Ser Leu Ala Ile Pro Cys Asn Asn Val Glu  
225 230 235 240

Val Leu Met Ser Gly Asn Val Arg Cys Tyr Asn Ile Val Val Glu Ala  
245 250 255

Met Lys Ala Phe Pro Met Ser Glu Arg Ile Gln Glu Val Ser Cys Cys  
260 265 270

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275 280 285

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305 310 315 320

Glu Thr Ile Phe Leu Asn Gln Asp Leu Glu Glu Lys Asn Glu Asn Gln  
325 330 335

Glu Asn Asp Asp Glu Gly Glu Glu Asp Lys Leu Phe Trp Leu Glu Ala  
340 345 350

Cys Tyr Lys Ala Leu Thr Trp His Arg Lys Asn Lys His Val Gln Glu  
355 360 365

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Lys Cys Gly Leu Lys Val Ile Ser Ser Ile Val His Phe Pro Asp Ala		
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Leu Glu Met Leu Ser Leu Glu Gly Ala Met Asp Ser Val Leu His Thr		
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Leu Ile Gly Tyr Leu Ile Thr Lys Lys Asn Val Phe Ile Gly Thr Gly		
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His Leu Leu Ala Lys Ile Leu Val Ser Ser Leu Tyr Arg Phe Lys Asp		
625	630	635
640		
Val Ala Glu Ile Gln Thr Lys Gly Phe Gln Thr Ile Leu Ala Ile Leu		
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Lys Leu Ser Ala Ser Phe Ser Lys Leu Leu Val His His Ser Phe Asp		
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770 775 780

Gln Ile Ile Ser Leu Leu Leu Arg Arg Leu Ala Leu Asp Val Ala Asn  
785 790 795 800

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805 810 815

Trp Leu Gly Pro Leu Phe Pro Asp Lys Thr Ser Asn Leu Arg Lys Gln  
820 825 830

Thr Asn Ile Ala Ser Thr Leu Ala Arg Met Val Ile Arg Tyr Gln Met  
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850 855 860

Ser Glu Asp Val Leu Ser Lys Phe Asp Glu Trp Thr Phe Ile Pro Asp  
865 870 875 880

Ser Ser Met Asp Ser Val Phe Ala Gln Ser Asp Asp Leu Asp Ser Glu  
885 890 895

Gly Ser Glu Gly Ser Phe Leu Val Lys Lys Lys Ser Asn Ser Ile Ser  
900 905 910

Val Gly Glu Phe Tyr Arg Asp Ala Val Leu Gln Arg Cys Ser Pro Asn  
915 920 925

Leu Gln Arg His Ser Asn Ser Leu Gly Pro Ile Phe Asp His Glu Asp  
930 935 940

Leu Leu Lys Arg Lys Arg Lys Ile Leu Ser Ser Asp Asp Ser Leu Arg  
945 950 955 960

Ser Ser Lys Leu Gln Ser His Met Arg His Ser Asp Ser Ile Ser Ser  
965 970 975

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980 985 990

Glu Leu Arg Asp Ile Asp Ala Leu Ser Gln Lys Cys Cys Ile Ser Val  
995 1000 1005

His Leu Glu His Leu Glu Lys Leu Glu Leu His Gln Asn Ala Leu  
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Thr Ser Phe Pro Gln Gln Leu Cys Glu Thr Leu Lys Ser Leu Thr  
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His Leu Asp Leu His Ser Asn Lys Phe Thr Ser Phe Pro Ser Tyr  
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Leu Leu Lys Met Ser Cys Ile Ala Asn Leu Asp Val Ser Arg Asn  
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Asp Ile Gly Pro Ser Val Val Leu Asp Pro Thr Val Lys Cys Pro  
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Thr Leu Lys Gln Phe Asn Leu Ser Tyr Asn Gln Leu Ser Phe Val  
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Val Gly Ile Asp Val Lys Asp Trp Pro Ile Gln Ile Arg Asp Lys  
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Arg Lys Arg Asp Leu Val Leu Asn Val Trp Asp Phe Ala Gly Arg  
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Asp Ala Met Lys Pro Trp Leu Phe Asn Ile Lys Ala Arg Ala Ser  
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Glu Lys Gln Arg Lys Ala Cys Met Ser Lys Ile Thr Lys Glu Leu  
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Ile Leu Ser Glu Arg Lys Asn Val Pro Ile Glu Phe Pro Val Ile  
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Asp Arg Lys Arg Leu Leu Gln Leu Val Arg Glu Asn Gln Leu Gln  
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Leu Asp Glu Asn Glu Leu Pro His Ala Val His Phe Leu Asn Glu  
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Ser Gly Val Leu Leu His Phe Gln Asp Pro Ala Leu Gln Leu Ser  
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1595 1600 1605

Gln Ile Leu Thr Val Lys Val Glu Gly Cys Pro Lys His Pro Lys  
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Glu Lys Phe Gln Ile Ala Leu Pro Ile Gly Glu Glu Tyr Leu Leu  
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Val Pro Ser Ser Leu Ser Asp His Arg Pro Val Ile Glu Leu Pro  
1670 1675 1680

His Cys Glu Asn Ser Glu Ile Ile Ile Arg Leu Tyr Glu Met Pro  
1685 1690 1695

Tyr Phe Pro Met Gly Phe Trp Ser Arg Leu Ile Asn Arg Leu Leu  
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Pro Asn Arg Met Tyr Trp Arg Gln Gly Ile Tyr Leu Asn Trp Ser  
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1745 1750 1755

Pro Glu Ser Phe Leu Lys Ile Thr Val Pro Ser Cys Arg Lys Gly  
1760 1765 1770

Cys Ile Leu Leu Gly Gln Val Val Asp His Ile Asp Ser Leu Met  
1775 1780 1785

Glu Glu Trp Phe Pro Gly Leu Leu Glu Ile Asp Ile Cys Gly Glu  
1790 1795 1800

Gly Glu Thr Leu Leu Lys Lys Trp Ala Leu Tyr Ser Phe Asn Asp  
1805 1810 1815

Gly Glu Glu His Gln Lys Ile Leu Leu Asp Asp Leu Met Lys Lys  
1820 1825 1830

Ala Glu Glu Gly Asp Leu Leu Val Asn Pro Asp Gln Pro Arg Leu  
1835 1840 1845

Thr Ile Pro Ile Ser Gln Ile Ala Pro Asp Leu Ile Leu Ala Asp  
1850 1855 1860

Leu Pro Arg Asn Ile Met Leu Asn Asn Asp Glu Leu Glu Phe Glu  
1865 1870 1875

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1880 1885 1890

Tyr Arg Ala Ala Tyr Glu Gly Glu Glu Val Ala Val Lys Ile Phe  
1895 1900 1905

Asn Lys His Thr Ser Leu Arg Leu Leu Arg Gln Glu Leu Val Val  
1910 1915 1920

Leu Cys His Leu His His Pro Ser Leu Ile Ser Leu Leu Ala Ala  
1925 1930 1935

Gly Ile Arg Pro Arg Met Leu Val Met Glu Leu Ala Ser Lys Gly  
1940 1945 1950

Ser Leu Asp Arg Leu Leu Gln Gln Asp Lys Ala Ser Leu Thr Arg  
1955 1960 1965

Thr Leu Gln